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<110> Brett P. Monia  
Kenneth W. Dobie

<120> ANTISENSE MODULATION OF HISTONE DEACETYLASE 2 EXPRESSION

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gtggcggcgg tggccgggga gccc atg gcg tac agt caa gga ggc ggc aaa 231

Met Ala Tyr Ser Gln Gly Gly Gly Lys

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aaa aaa gtc tgc tac tac tac gac ggt gat att gga aat tat tat tat 279

Lys Lys Val Cys Tyr Tyr Tyr Asp Gly Asp Ile Gly Asn Tyr Tyr Tyr

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Gly Gln Gly His Pro Met Lys Pro His Arg Ile Arg Met Thr His Asn

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ttg ctg tta aat tat ggc tta tac aga aaa atg gaa ata tat agg ccc 375

Leu Leu Leu Asn Tyr Gly Leu Tyr Arg Lys Met Glu Ile Tyr Arg Pro

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cat aaa gcc act gcc gaa gaa atg aca aaa tat cac agt gat gag tat 423

His Lys Ala Thr Ala Glu Glu Met Thr Lys Tyr His Ser Asp Glu Tyr

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Ile Lys Phe Leu Arg Ser Ile Arg Pro Asp Asn Met Ser Glu Tyr Ser	
75 80 85	
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Lys Gln Met His Ile Phe Asn Val Gly Glu Asp Cys Pro Ala Phe Asp	
90 95 100 105	
gga ctc ttt gag ttt tgt cag ctc tca act ggc ggt tca gtt gct gga	567
Gly Leu Phe Glu Phe Cys Gln Leu Ser Thr Gly Gly Ser Val Ala Gly	
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Ala Val Lys Leu Asn Arg Gln Gln Thr Asp Met Ala Val Asn Trp Ala	
125 130 135	
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Gly Gly Leu His His Ala Lys Lys Tyr Glu Ala Ser Gly Phe Cys Tyr	
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Val Asn Asp Ile Val Leu Ala Ile Leu Glu Leu Lys Tyr His Gln	
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Arg Val Leu Tyr Ile Asp Ile Asp Ile His His Gly Asp Gly Val Glu	
170 175 180 185	
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Glu Ala Phe Tyr Thr Thr Asp Arg Val Met Thr Val Ser Phe His Lys	
190 195 200	
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Tyr Gly Glu Tyr Phe Pro Gly Thr Gly Asp Leu Arg Asp Ile Gly Ala	
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Gly Lys Gly Lys Tyr Tyr Ala Val Asn Phe Pro Met Cys Asp Gly Ile	
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Asp Asp Glu Ser Tyr Gly Gln Ile Phe Lys Pro Ile Ile Ser Lys Val	
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Thr Tyr Glu Thr Ala Val Ala Leu Asp Cys Glu Ile Pro Asn Glu Leu	
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Pro Tyr Asn Asp Tyr Phe Glu Tyr Phe Gly Pro Asp Phe Lys Leu His	
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cct ggt gtc cag atg caa gct att cca gaa gat gct gtt cat gaa gac	1383
Pro Gly Val Gln Met Gln Ala Ile Pro Glu Asp Ala Val His Glu Asp	
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Ser Gly Asp Glu Asp Gly Glu Asp Pro Asp Lys Arg Ile Ser Ile Arg	
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Ala Ser Asp Lys Arg Ile Ala Cys Asp Glu Glu Phe Ser Asp Ser Glu	
410	415 420 425
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Asp Glu Gly Glu Gly Gly Arg Arg Asn Val Ala Asp His Lys Lys Gly	

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Ala Lys Lys Ala Arg Ile Glu Glu Asp Lys Lys Glu Thr Glu Asp Lys			
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Lys Thr Asp Val Lys Glu Glu Asp Lys Ser Lys Asp Asn Ser Gly Glu			
460	465	470	
aaa aca gat acc aaa gga acc aaa tca gaa cag ctc agc aac ccc tga			1671
Lys Thr Asp Thr Lys Gly Thr Lys Ser Glu Gln Leu Ser Asn Pro			
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PTS-0022

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